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May 19, 2003, 16:36:02 ; Search time 15.2044 Seconds (without alignments) 982.052 Million cell updates/sec
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1900
T-MLSTSRSFRFIRNTNESGEEV.....DGVTSTNTPSTGEQEVSAGL 360
GenCore version 5.1.4\_p5\_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                            112892 seqs, 41476328 residues
                                                                              OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Perfect score:
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112892 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 08 Maximum Match 1008 Listing first 45 summaries

SwissProt 40:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

No. Score Match Length DB ID  1 1845 97.1 360 1 CKR2_MACMU 3 1542 81.2 373 1 CKR2_RAT 4 1535 80.8 373 1 CKR2_RAT 5 1396 72.9 354 1 CKR2_MOUSE 6 1386 72.9 354 1 CKR5_RAT 7 1376 72.4 352 1 CKR5_RAT 10 1370 72.1 352 1 CKR5_RAT 11 1370 72.1 352 1 CKR5_GNGO 11 1370 72.1 352 1 CKR5_GNGO 11 1370 72.1 352 1 CKR5_GNRO 11 1370 72.1 352 1 CKR5_GNRO 11 1370 72.1 352 1 CKR5_GNRO 11 1368 72.0 352 1 CKR5_GNRO 11 1368 72.0 352 1 CKR5_GNRO 12 1368 72.0 352 1 CKR5_GNRO 13 1368 72.0 352 1 CKR5_PAPHA 15 1363 71.7 352 1 CKR5_PAPHA 15 1363 71.7 352 1 CKR5_PAPHA 15 1364 71.7 352 1 CKR5_PYONE 16 1361 71.6 352 1 CKR5_PYONE 17 1360 71.6 352 1 CKR5_PYONE 18 1359 71.5 352 1 CKR5_PYONE 19 1344 70.7 352 1 CKR5_PYONE 22 1003.5 52.8 359 1 CKR3_MOUSE 24 980 51.6 355 1 CKR3_MOUSE 25 940.5 50.6 355 1 CKR3_MOUSE 26 947 49.8 355 1 CKR3_MOUSE 27 940.5 49.7 355 1 CKR3_MOUSE 28 940.5 49.7 355 1 CKR3_MOUSE 29 94.5 48.7 355 1 CKR4_MOUSE 29 94.5 48.7 355 1 CKR4_MOUSE 21 778.5 41.0 356 1 CKR8_MOUSE 22 778.5 39.1 CKR8_MACMU 23 743.5 39.1 CKR8_MACMU 24 981 74.5 48.7 355 1 CKR8_MACMU 25 778.5 39.1 CKR8_MACMU			æ			SUMMARIES		
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1863   71.7   352   CKR   1360   71.6   352   CKR   1360   71.6   352   CKR   2344   70.7   352   CKR   2344   70.7   352   CKR   2344   70.7   352   CKR   235	14	1364		352	Н	CKR5_HUMAN	P51681	homo sapien
7 1361 71.6 352 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	15	1363	71.	352	Н	CKR5_TRAPH	628260	trachypithe
7 1360 71.6 352 1 33 1344 70.7 352 1 134 70.7 352 1 1018.5 53.6 359 1 2 1003.5 52.8 359 1 2 1003.5 52.8 359 1 4 981 51.6 359 1 5 940.5 50.6 355 1 6 940.7 49.8 355 1 8 940.5 48.7 355 1 8 940.5 48.7 355 1 8 74.5 48.7 355 1 8 778.5 41.0 356 1 778.5 41.0 356 1 778.5 39.1 356 1	16	1361	71.	352		CKR5_PYGBI	097880	pygathrix b
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1 778.5 41.0 356 1 2 751 39.5 355 1 3 743.5 39.1 356 1	30	862.5	45	360		CKR4_HUMAN	P51679	homo sapien
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3 743.5 39.1 356 1	32	5		355	~	CKR8_HUMAN	P51685	homo sapien
	33	43.	39.1	356	7	CKR8_MACMU	097665	macaca mula

P56484 mus musculu P35411 rattus norv O92409 mus musculu P49238 homo sapien 006590 homo sapien 008907 rattus norv P32248 homo sapien P51686 homo sapien O54689 mus musculu Q9wut7 mus musculu P47774 mus musculu		SSULT 1  CRR2_MACMU CRR2_MACMU STANDARD; PRT; 360 AA.  CORP3_3.  16-OCT-2001 (Rel. 40, Last sequence update)  F 16-OCT-2001 (Rel. 40, Last annotation update)  C C chemokine receptor type 2 (C-C CRR-2) (CCR-2) (CCR2)  C C chemokine receptor protein 1 receptor) (MCP-1-R) (CCR2).	Vertebrata; Euteleostomi; i; Cercopithecidae;	J.E.; even rhesus macaque chemokine	((2001)). MCP-3 AND MCP-4 CHEMOKINES. THE INTRACELLULAR CALCIUM IONS	LEVEL. SUBCELLULAR LOCATION: Integral membrane protein. ALTERNATIVE PRODUCTS: 2 ISOFORMS; A AND B (SHOWN HERE); ARE PRODUCED BY ALTERNATIVE SPLICING. SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.	is produced through a collaboration rmatics and the EMBL outstation - There are no restrictions on its nog as its content is in no way weed. Usage by and for commercial see http://www.isb-sib.ch/announce/	1. 1. nne; Glycoprotein; Sulfation; HAR (POTEWHIAL).	FT TRANSMEM 43 70 1 (POTENTIAL).  FT DOMAIN 71 80 CYTOPLASMIC (POTENTIAL).  FT DOMAIN 101 114 EXTRACELLULAR (POTENTIAL).  FT TRANSMEM 115 136 3 (POTENTIAL).  FT TRANSMEM 137 153 CYTOPLASMIC (POTENTIAL).  FT TRANSMEM 154 178 4 (POTENTIAL).  FT TRANSMEM 179 206 EXTRACELLULAR (POTENTIAL).  FT TRANSMEM 207 226 5 (POTENTIAL).
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MEDLINE-97150864; Pubmed-8995400;
Wong L.-M., Myers S.J., Tsou C.-L., Gosling J., Arai H., Charo I.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine receptor type 2 (C-C CKR-2) (CC-CKR-2) (CCR2)
(Monocyte chemoattractant protein 1 receptor) (MCP-1-R) (CCR2).
                                                                                                                                                                                                                                                                                                                                             CQKEDSVYVCGPYFPRGWNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Molecular cloning and functional expression of two monocyte chemoattractant protein 1 receptors reveals alternative splicing the carboxyl-terminal tails.";
                                                                    (POTENTIAL)
                                                                                                                                                               ö
                                                                                                                                 Score 1845; DB 1; Length 360;
Pred. No. 1.3e-101;
5; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Charo I.F., Myers S.J., Herman A., Franci C., Connolly A.J., Coughlin S.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yamagami S., Tokuda Y., Ishii K., Tamaka H., Endo N.; "cDNA cloning and functional expression of a human monocyte chemoattractant protein 1 receptor."; Biochem. Biophys. Res. Commun. 202:1156-1162(1994).
               6 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
                                       7 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
N-LINKED (GLCNAC. . ) (PO'SULFATION (BY SIMILARITY).
BY SIMILARITY.
                                                                                                          4B2552BCE913FE9F CRC64;
    CYTOPLASMIC (POTENTIAL).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            374 AA.
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MEDLINE-94195821; PubMed-8146186;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDLINE=94324942; PubMed=8048929;
                                                                                                        41139 MW;
                                                                                                                                 97.18;
96.98;
                                                                                                                                                          Matches 349; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
 243
268
285
309
360
14
26
                                                                                                       360 AA;
                                                                                                                                             Similarity
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CKR2_HUMAN
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                                                                                                                                                                                                                                                           MEDLINE-2001139; PubMed-11046064; Preobrazhensky A.A., Dragan S., Kawano T., Gavrilin M.A., Gulina I.V., Chakravarty L., Kolattukudy P.E.; "Monocyte chemotactic protein-1 receptor CCR2B is a glycoprotein that has tyrosine sulfation in a conserved extracellular N-terminal
"Organization and differential expression of the human monocyte chemoattractent protein 1 receptor gene. Evidence for the role of the carboxyl-terminal tail in receptor trafficking."; J. Biol. Chem. 272:1038-1045(1997).
                                                                                                                                                                                                                                                                                                                                                           J. Immunol. 165:5295-5303(2000).
-!- FUNCTION: Receptor for the MCP-1, MCP-3 and MCP-4 chemokines.
Transduces a signal by increasing the intracellular calcium ions level. Alternative coreceptor with CD4 for HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -!- PTM: N-GLYCOSYLATED.
-!- SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
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PROSITE; PS50262; G_PROTEIN_RECEP_FI_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation; Alternative splicing.
                                                                                              McCombie W.R., Wilson R., Chen E., Gibbs R., Zuo L., Johnson D., Nhan M., Parnell L., Dedhia N., Ansari A., Mardis E., Schutz K., Gnoj L., la Bastide M., Kaplan N., Grecor T., Touchman J., Muzny D., Chen C.N., Evans C., Fitzgerald M., See L.H., Tang M., Forcel B.M., Dragan Y., Giacalone J., Pae A., Powell E., Solinsky K.A., Desilva U., biaz-Perez S., Zhou X., Yu Y., Watanabe M., Doggett N., Garcia D., Sagripanti J.L.; Submitted (MAY-1997) to the EMBL/GenBank/DDBJ databases.
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CYTOPLASMIC (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                            -!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- ALTERNATIVE PRODUCTS: 2 ISOFORMS; A (SHOWN HERE) AND B; ARE PRODUCED BY ALTERNATIVE SPLICING.
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EXTRACELLULAR (POTENTIAL).
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3 (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
6 (POTENTIAL).
EXTRECELLULAR (POTENTIAL).
7 (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
2 (POTENTIAL).
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                                                                                                                                                                                                                                [5] SULFATION OF TYR-26, AND N-GLYCOSYLATION.
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EMBL, U03905; AAA19120.1; --
EMBL, D29984; BAA06253.1; --
EMBL, U80924; AAC51637.1; --
EMBL, U95626; AAB57791.1; --
EMBL, U95626; AAB57791.1; --
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1114
1136
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2226
2243
374
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                                                                                   SEQUENCE FROM N.A.
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101
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J. Neuroimmunol. 86:1-12(1998).
-!- FUNCTION: RECEPTOR FOR THE MOP-1 (JE), MCP-3 (FIC) AND MCP-5 CHEMOKINES. TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLUIAR CALCIUM IONS LEVEL (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- TISSUE SPECIFICITY: EXPRESSED IN LUNG, SPLEEN, KIDNEY, THYMUS AND
BY SIMILARITY
SLFHIALGCRIAPLOKPVCGGPGVRPGKNVKVTTOGLLDGR
                                         RYLSVFFRKHITKRFCK
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                                                 1 MLSTSRSRFIRNTNESGEEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGN 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
C-C chemokine receptor type 2 (C-C CKR-2) (CCR-2) (CCR2)
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                                 GKGKSIGRAPEASLQDKEGA -> RYLSVFFRKHI
QCPVFYRETVDGVTSTNTPSTGEQEVSAGL (IN
                                                                                                                           DB 1; Length 374;
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Pred. No. 2.7e-90;
....rrrhes 5; Indels
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                                                                       ISOFORM B)
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MEDLINE-98318173; PubMed-9655467;
                                                                                           MW.
                                                                                                                           86.98;
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Conservative
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Rattus norvegicus (Rat).
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30-MAY-2000 (Rel. 39, Last annotation update)
C-C chemokine receptor type 2 (C-C CKR-2) (CC-CKR-2) (CCR-2) (CCR-2)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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EXTRACELLULAR (POTENTIAL).
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EXTRACELLULAR (POTENTIAL)
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80.0%; Pred. No. 6.8e-84;
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                                                               EMBL; U77349; AAC03242.1; -.
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
PR00179; PR00237; GPCRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BY SIMILARITY
                                                                                                                                                                                                               G-protein coupled receptor; Transmembrane.
entitles requires a license agreement (or send an email to license@isb-sib.ch)
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CKR2_MOUSE Sich...
P51683: Q61172:
01-OCT-1996 (Rel. 34, Created)
...nv-1997 (Rel. 35, Last seque
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SEQUENCE FROM N.A.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                   chemoattractant protein 1) and murine macrophage inflammatory protein
lalpha receptors: evidence for two closely linked C-C chemokine
                                                                                                                                                                                                                                                                                                    MEDLINE=97026720; PubMed=8872898; Heesen M., Tanabe S., Berman M.A., Yoshizawa I., Luo Y., Kim R., Heesen M., Tanabe S., Berman M.A., Yoshizawa I., Luo Y., Kim R., Post T.W., Gerard C., Dorf M.E.; Mouse astrocytes respond to the chemokines MCP-1 and KC, but reverse transcriptase-polymerase chain reaction does not detect mRNA for the KC or new MCP-1 receptor.";
                                                                                                                                                                                                                                                                                                                                                                                                      J. Neurosci. Res. 45:882-391(1996).
-!- FUNCTION: RECEPTOR FOR THE MCP-1 (JE), MCP-3 (FIC) AND MCP-5 CHEMOKINES. TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL.
-!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- TISSUE SPECIFICITY: DETECTED IN MONOCYTE/MACROPHAGE CELL LINES, BUT NOY IN NONHEMATPOLETIC CELL LINES.
-!- SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                 Molecular cloning and functional expression of murine JE (monocyte
                                                                                                                                                                                                      Kurihara T., Bravo R.;
"Cloning and functional expression of mCCR2, a murine receptor for
              Boring L., Gosling J., Monteclaro F.S., Lusis A.J., Tsou C.-L., Charo I.F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
2 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
4 (POTENTIAL).
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5 (POTENTIAL).
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EXTRACELLUIAR (POTENTIAL).
7 (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
BY SIMILARITY.
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-> G (IN REF. 1).
-> G (IN REF. 1).
FA012C10F4C9325A CRC64;
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PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
G-PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-Protein coupled receptor; Transmembrane.
                                                                                                                                                                                                                                     the C-C chemokines JE and FIC.";
J. Biol. Chem. 271:11603-11606(1996).
                                                                                                 receptors on chromosome 9.";
J. Biol. Chem. 271:7551-7558(1996).
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InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                   STRAIN=BALB/c;
MEDLINE=96216064; PubMed=8662823;
 MEDLINE=96205938; PubMed=8631787;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; U47035; AAC52453.1; -. EMBL; U51717; AAC52557.1; -. EMBL; U56819; AAC52784.1; -.
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264
373 AA;
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Length 373;

DB 1;

Score 1535;

Query Match

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"Molecular cloning and functional expression of murine JE (monocyte chemoattractant protein 1) and murine macrophage inflammatory protein lalpha receptors: evidence for two closely linked C-C chemokine
                                                                                                                                                                            74 MLVIILLIGCKKLKSMTDIYLLNLAISDLLFLLTLPFWAHYAANEWVFGNIMCKVFTGLY 133
                                                                                                                                                                                                                                                       181 CQKEDSVYVCGPYFPRGWNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHR 240
                                                                                                                                                                                                                                                                                                                                61 MLVVLILLINCKKLKCLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLY 120
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                                                                                               14 ILSTSHSLFTRSIQELDEGATTPYDYDDGEPCHKTSVKQIGAWILPPLYSLVFIFGFVGN 73
                                                                           1 MLSTSRSRFIRNTNESGEEVTTFFDYDYGAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGN 60
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P51682; Q61867; P97405; O35313; P97308; O35891;
01-0CT-1996 (Rel. 34, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (MIP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=C57BL/6, and NIH Swiss; TISSUE=Liver, Kidney, and Spleen; MEDLINE=98001387; PubMed=9341222; Kuhmann S.E., Platt E.J., Kozak S.L., Kabat D.; Kuhmann S.E., Polymorphisms in the CCR5 genes of African green monkeys and mice implicate specific amino acids in infections by simian and human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=C57BL/6 X CBA; TISSUE=Thymus;
MEDLINE=96278910; PubMed=8662890;
Meyer A., Coyle A.J., Proudfoot A.E.I., Wells T.N.C., Power C.A.;
"Cloning and characterization of a novel murine macrophage
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MEDLINE=96205938; PubMed=8631787;
Boring L., Gosling J., Monteclaro F.S., Lusis A.J., Tsou C.-L.,
                                      Indels
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Kuziel W.A., Beck M.A., Dawson T.C., Maeda N.;
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
                                  48;
                    Pred. No. 1.7e-83;
                                        Mismatches
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J. Biol. Chem. 271:14445-14451(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Biol. Chem. 271:7551-7558(1996).
79.78; Fr. 25;
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                                    Matches 287; Conservative
                  Best Local Similarity
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CCR5 OR CMKBR5.
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                                                                                                                                                                                                                                                                                                                                                                                    Guo B., Kuno K., Harada A., Matsushima K.;
Submitted (JAN-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY
INCREASING THE INTRACELLUIAR CALCIUM IONS LEVEL.
                                                            Doranz B.J., Lu Z.H., Rucker J., Zhang T.Y., Sharron M., Cen Y.H., Wang Z.X., Guo H.H., Du J.G., Accavitti M.A., Doms R.W., Peiper S.C., "Two distinct CCR5 domains can mediate coreceptor usage by human immunodeficiency virus type 1";
J. Virol. 71:6305-6314(1997).
                                                                                                                                                                                    SUBCELLUIAR LOCATION: Integral membrane protein.
TISSUE SPECIFICITY: DETECTED IN MONOCYTE/MACROPHAGE CELL LINES,
BUT NOT IN NONHEMATOPOIETIC CELL LINES.
SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
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EXTRACELLULAR (POTENTIAL).

7 (POTENTIAL).
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          Virol. 71:8642-8656(1997).
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EMBL; X94151; CAA63867.1; --
EMBL; U68565; AAB37273.1; --
EMBL; U83327; AAC53386.1; --
  immunodeficiency viruses.";
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160
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EMBL;
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STRAIN=Sprague-Dawley;
MEDLIND=98318173; PubMed=9655467;
Jiang Y., Salafranca M.N., Adhikari S., Xia Y., Feng L., Sonntag M.K.,
Defiebre C.M., Pennell N.A., Streit W.J., Harrison J.K.;
"Chemokine receptor expression in cultured glia and rat experimental
                                                                                                                                                                                                                                                                                                                                                                                                310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Neuroimmunol. 86:1-12(1998).
-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA, MIP-1-BETA AND KANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL.
-!- SUBCELLULAR LOCATION: Integral membrane protein.
-!- SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                                                                                                                                                                                75 CLTDIYLLNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILL 134
                                                                                                                                                                                                                                                                                          135 TIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYF 194
                                                                                                                                  Gaps
                                                                                                                                                       Eukaryota; Metazoa; Chórdata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBL_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-NOV-1997 (Rel. 35, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (MIP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Spleiss O., Gourmala N., Boddeke H.W.G.M., Sauter A., Fiebich B.L., Berger M., Gebicke-Haerter P.J.;
"Cloning of rat HIV-1-chemokine coreceptor CKR5 from microglia and upregulation of its mRNA in ischemic and endotoxinemic rat brain.";
J. Neurosci. Res. 53:16-28(1998).
                                                                                                                                                                                                                                                                                                                                                                                          VYFLFWTPYNIVILLNTFQEFFGLSNCESTSQLDQATQVTETLGMTHCCINPIIYAFVGE
                                                                                                                                                                                                                                                                                                                                   PRG----WNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMI
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9
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F -> L (IN REF. 2).
L -> F (IN REF. 2).
N -> I (IN REF. 5).
H -> Y (IN REF. 3).
P -> S (IN REF. 1).
W; B4A6B942E88F9CFO CRC64;
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                                                                                                  Score 1396;
Pred. No. 2.
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MEDLINE=98334064; PubMed=9670989;
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01-NOV-1997 (Rel. 35, Last sequ
30-MAY-2000 (Rel. 39, Last anno
                                                                     40863 MW;
                                                                                                  73.58;
74.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               allergic encephalomyelitis."
                                                                                                            Local Similarity, 74.6
nes 261; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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3
80
145
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208
                                                                    354 AA;
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008556:
                                                                     SEQUENCE
                                                                                                  Query Match
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                        This SWISS-PROT entry is copyright. It is produced through a collaboration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       195 ----PRGWNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMI 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   135 TIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCOKEDSVYVCGPYF 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                184
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062743; 062744; 062745; 062746;
15-DEC-1998 (Rel. 37, Last sequence update)
115-DEC-1998 (Rel. 40, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CCR-5) (CCR5).
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                                                                                                                                                                                                                                                                                                          Pfam; PF00001; /tm__i, __.
PR0137; PF0RRHODPSN.
PROSITE: PS0237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE: PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein.
32 EXTRACELLULAR (POTENTIAL).
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EXTRACELLULAR (POTENTIAL).
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EXTRACELLUAR (POTENTIAL).
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EXTRACELLULAR (POTENTIAL).
5 (POTENTIAL).
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Pred. No. 8.6e-75;
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Pfam; PF00001; 7tm_1; 1.
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74.98;
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                                                                                                                                                                         naturally infected in west Africa: a comparison of coreceptor usage of primary SIVSm, HIV-2, and SIVmac.";
Virology 246:113-124(1998).

- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA, MIP-1-BETTA AND RAWIES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEE. MAY PLAY A ROLE IN THE CONTROL OF GRANULCCYTIC LINEAGE PROLIFERATION OR
                                                                                                               STRAIN=Isolate 079, 085, 087, and 089;
MEDLINE=98321155; PubMed=9656999;
Chen Z., Gettie A., Ho D.D., Marx P.A.;
"Primary SIVsm isolates use the CCR5 coreceptor from sooty mangabeys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00237; CMTRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.
                                                                                                                                                                                                                                                                                             -i- SUBCELLULAR LOCATION: Integral membrane protein.
             Cercocebus torquatus atys (Red-crowned mangabey) (Sooty mangabey). Bukaryota Metazoa: Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria: Primates; Catarrhini; Cercopithecidae; Cercocbus.
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SULFATION (BY SIMILARITY).
D -> E (IN ISOLATE 087).
Y -> D (IN ISOLATE 079).
V -> G (IN ISOLATE 079).
M -> K (IN ISOLATE 089).
L -> V (IN ISOLATE 089).
V -> G (IN ISOLATE 089).
V -> G (IN ISOLATE 079).
T -> I (IN ISOLATE 079).
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CYTOPLASMIC (POTENTIAL).
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EXTRACELLUIAR (POTENTIAL).
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20A196E2D47E49CA CRC64;
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Pred. No. 3.3e-74;
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EMBL, AF051903, AAC39831.1; -.
EMBL, AF051904, AAC39832.1; -.
EMBL, AF051905, AAC39833.1; -.
InterPro, IPR0000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1
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Best Local Similarity
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                                                                                                    SEQUENCE FROM N.A.
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59
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90
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CCR5 OR CMKBR5.
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Zhang Y.-W., Ryder O.A., Zhang Y.-P.;
Zhang Y.-W., Ryder O.A., Zhang Y.-P.;
Sequence evolution of the CCRS chemokine receptor gene in primates.";
Mol. Biol. Evol. 16:1145-1154(1999).
-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY
INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL. MAY PLAY A ROLE
IN THE CONTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                                                                                                      LNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA 141
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   Gaps
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15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCRCS).
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PRINTS; PRO0237; GPRRHODOPSN.

PROSITE; PS00237; G_PROTEIN.RECEP_FI_1; 1.

PROSITE; PS50262; G_PROTEIN_RECEP_FI_2; 1.

G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.

DOMAIN 1 30 EXTRACELLULAR (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                  FDYDY -- GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL
                                                      PYNIVILLNTFQEFFGLSNCESTSQLDQATQVTETLGMTHCCINPIIYAFVGEKFRRYLS
                                                                                                                                                                                                                                          WNNFHTIMRNILGLVLPLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hylobates leucogenys (White-cheeked gibbon).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hylobatidae; Hylobates.
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9
     44; Indels
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CYTOPLASMIC (POTENTIAL).
2 (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                   VFFRKHITKRFCKQCPVFYRETVDGVTSTNTPSTGEQEVSAGL 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         352 AA
   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-MAY-2000 (Rel. 39, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AF075451; AAD19863.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=61853;
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097883;
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 Matches
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312
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                                                                                                                                                                                                                                                                             AISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLAIVH 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CKR5_MACMU STANDARD; PRT; 352 AA.
P79436; 002746;
01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CCR-5) (CCR5).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCR5 OK CMRDKO.
Macaca mulatta (Rhèsus macaque),
Macaca mulatta (Rhèsus macaque),
Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey), and
Macaca nemestrina (Pig-tailed macaque).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                 DYDYGAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYLLNL
                                                                                                                                                                                                                                               AVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RGWNN
                                                                                                                                                                                                                                                                                                                                                                     201 FHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWTPYN
                                                                                                                                                                                                                                                                                                                                                                               L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-97184592; PubMed-9032394; Marcon L., Choe H., Martin K.A., Farzan M., Ponath P.D., Wu Newman W., Gerard N., Gerard C., Sodroski J.; Wu Tillization of C-C chemokine receptor 5 by the envelope 91ycoproteins of a pathogenic simian immunodeficiency virus, SIVMac239.";
                                                                                                                                                                                         DB 1; Length 352;
          3 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
4 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
                                                                                                                    BY SIMILARITY.
SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
AFBRAF344CEB7C91 CRCG4,
  EXTRACELLULAR (POTENTIAL).
                                                                          6 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
7 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
                                            5 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
6 (POTENTIAL).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 1371; DB 1,
                                                                                                                                                                                                  Pred. No. 6.46
                                                                                                                                                                  MM:
                                                                                                                                                                                      72.2%;
76.2%;
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                                                                                                                                                                 40445
                                                                                                                                                                                                            Conservative
102
1124
1141
1166
1166
235
235
2260
227
3301
3352
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                                                                                                                                                                352 AA;
                                                                                                                                                                                                 Similarity
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90
103
1125
1142
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1199
2219
2219
2261
2278
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DOMAIN
TRANSMEM
                                                                                                            DOMAIN
DISULFID
                                                                                                                                                      MOD_RES
SEQUENCE
                                                                          TRANSMEM
                                 PRANSMEM
                                                    FRANSMEM
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                    DOMAIN
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                     SPECIES—M. M. fascicularis, and M. nemestrina;
MEDLINE-97268687; Pubbed-9108095;
MEDLINE-97268687; Pubbed-9108095;
Beinger A.L., Amedee A., Miller K., Doranz B.J., Endres M.,
Sharron M., Lu Z.H., Clements J.E., Murphey-Corb M.,
Peiper S.C., Parmentier M., Broder C.C., Doms R.W.;
"Differential utilization of CCR5 by macrophage and T cell tropic
simian immunodeficiency virus strains.";
Proc. Natl. Acad. Sci. U.S.A. 94.4005-4010(1997).
- FUNCTION: RECEPTOR FOR A C. TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
MIP-1-BETA AND RAWYES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY
INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL. MAY PLAY A ROLE
SEQUENCE FROM N.A.
SPECIESAM mulatta; STRAIN-Indian macaque;
MEDINE-9721394; PubMed=9060623;
Chen Z., Zhou P., Ho D.D., Landau N.R., Marx P.A.;
Generically divergent strains of simian immunodeficiency virus use
"Geneptor for entry.";
J. Virol. 71:2705-2714(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIMITS, PRO0237, GENOTEIN RECEP_FI_1; 1.
PROSITE; PS00262; G_PROTEIN_RECEP_FI_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.
DOMAIN 1 30 EXTRACELLUIAR (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                                                                                                                                            Margulies B.J., Hauer D.A., Clements J.E.; "Identification and comparison of eleven rhesus macague chemokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                       IN THE CONTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
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CYTOPLASMIC (POTENTIAL).
4 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
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EXTRACELLULAR (POTENTIAL)
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CYTOPLASMIC (POTENTIAL).
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                                                                                                                                                                                                                             AIDS Res. Hum. Retroviruses 17:981-986(2001)
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Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                 MEDLINE=21354176; PubMed=11461684;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL, U77672; AAC51109.1; -. EMBL, U73739; AAC51158.1; -. EMBL; U96762; AAC34132.1; -.
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68
89
102
124
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352
178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DIFFERENTIATION.
                                                                                                                                              SPECIES=M.mulatta;
                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                               SEQUENCE FROM N.A.
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236
261
261
278
302
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DOMAIN
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TRANSMEM
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TRANSMEM
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142 IVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RG 197
                                                                                                                                                                                          LNLAISDLEFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA 141
                                                                                                                                                                                                                                                                                                                                                                        PYNIVILLNTFQEFFGLSNCESTSQLDQATQVTETLGMTHCCINPIIYAFVGEKFRRYLS 317
                                                                                                      Gaps
                                                                                                                                              15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-Cochemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCR-5).
                                                                                                                                24 FDYDY -- GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-97426118; PubMed-9282822;
Zacharova V., Zachar V., Goustin A.S.;
"Sequence of chemokine receptor gene CCR5 in chimpanzees, a natural
HIV type 1 host.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.

MEDLINE=97268687; PubMed=9108095;
Edinger A.L., Amedee A., Miller K., Doranz B.J., Endres M.,
Sharron M., Samson M., Lu Z.H., Clements J.E., Murphey-Corb M.,
Peiper S.C., Parmentier M., Broder C.C., Doms R.W.;
"Differential utilization of CCR5 by macrophage and T cell tropic similan immunodeficiency virus strains.";
Proc. Natl. Acad. Sci. U.S.A. 94:4005-4010(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zhang L., Carruthers C.D., He T., Huang Y., Cao Y., Wang G., Hahn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pan troglodytes (Chimpanzee).
Sukaryota: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Pan.
NCBL_TaxID=9598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "HIV type 1 subtypes, coreceptor usage, and CCR5 polymorphism."; AIDS Res. Hum. Retroviruses 13:1357-1366(1997).
.) (POTENTIAL).
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9
                                                                     Length 352;
                                                                                                   45; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zimmerman P.A., Buckler-White A., Alkhatib G.;
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
          -> I (IN REF. 3).
-> M (IN REF. 3).
58B96C85909FACB2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                   318 VFFRKHITKRFCKQCPVFYRETVDGVTSTNTPSTGEQEVSAGL 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                    310 VFFQKHIAKRFCKCCSIFQQEAPERASSVYTRSTGEQEISVGL 352
                                                                     Score 1370; DB 1;
Pred. No. 7.4e-74;
N-LINKED (GLCNAC.
                                                                                                      Mismatches
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                                                                                                   31;
                                           40507 MW;
                                                                       72.18;
76.18;
                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
268
241
292
352 AA;
                                                                                       Similarity
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                                                                       Query Match
Best Local Simi
Matches 261;
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            CONFLICT
CONFLICT
SEQUENCE
 CARBOHYD
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130 IVHAVFALKARTVTFGVVTSVITWVVAVFASLPGIIFTRSQKEGLHYTCSSHFPYSQYQF 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                           PRT;
                                                                                                                                                                                              30-MAY-2000 (Rel. 39, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AF075446; AAD19858.1;
                                                                                                                                                                                                                                                Pongo pygmaeus (Orangutan)
                                                                                                                                                                           STANDARD;
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89
102
1102
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1198
2315
235
235
237
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3352
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                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                              NCBI_TaxID=9600;
                                                                                                                                                                                                                                       CCR5 OR CMKBR5.
                                                                                                                                                                           CKR5_PONPY 097881;
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DOMAIN
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                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
                       SEQUENCE FROM N.A.

MEDLINE=98090115; PubMed=9430250;

MEDLINE=98090115; PubMed=9430250;

Pretet J.-L., Zerbib A., Girard M., Guillet J.-G., Butor C.;

"Chimpanzee CXCR4 and CCR5 act as coreceptors for HIV type 1.";

AIDS Res. Hum. Retroviruses 13:1583-1587(1997).

-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,

MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY

INCREASING THE INTRRACELLULAR CALCIUM IONS LEVEE. MAY PLAY A ROLE

IN THE CONTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           142 IVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RG 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82 LNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA 141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24 FDYDY --GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL 81
                                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: Integral membrane protein. SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
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CYTOPLASMIC (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
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   AIDS Res. Hum. Retroviruses 13:1159-1161(1997).
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                [5]
SEQUENCE
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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Zhang Y.-W., Ryder O.A., Zhang Y.-P.;
Zhang Y.-W., Ryder O.A., Zhang Y.-P.;
Sequence evolution of the CCR5 chemokine receptor gene in primates.";
Mol. Biol. Evol. 16:1145-1154(1999).
-i- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
MIP-1-BETA AND RAWRES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY
INCREASING THE INTRACELLIARA CALCIUM IONS LEWEL. MAY PLAY A ROLE
IN THE CONTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                   258 PYNIVILLINTFOEFFGLSNCESTSQLDQATQVTETLGMTHCCINPIIYAFVGEKFRRYLS 317
WNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWT 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCR-5).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS00262; G_PROTEIN_RECEP_F1_2; 1.
G_PROTEIN_RECEP_F1_2; 1.
DOMAIN 1 30
FETEBACETITIES (A)ycoprotein; Sulfation.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pongo.
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CYTOPLASMIC (POTENTIAL).
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BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-97268687; PubMed=9108095; Edinger A.L., Amedede A., Miller K., Doranz B.J., Endres M., Sharron M., Samson M., Lu Z.H., Clements J.E., Murphey-Corb M., Peiper S.C., Parmentier M., Broder C.C., Doms R.W., "Differential utilization of CCR5 by macrophage and T cell tropic simian immunodeficiency virus strains ". Proc. Natl. Acad. Sci. U.S.A. 94:4005-4010(1997).

-I-FUNCTION: RECEPTOR FOR A C.-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA, MIP-1-BETA AND RAWIES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL. MAY PLAY A ROLE INTRACOUTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                                                                                                                                                                                                                                          82 LNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA 141
                                                                                                                                                                                                                                                                                                        142 IVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RG 197
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                                                                                                                                                                                                                                                                                                                                                                                              258 PYNIVILLNTFQEFFGLSNCESTSQLDQATQVTETLGMTHCCINPIIYAFVGEKFRYLS 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                     24 FDYDY -- GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL 81
                                                                                                                                                                                            15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
C-C chemokine receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCRCR5).
CCR5 OR CMKBR5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBCELLULAR LOCATION: Integral membrane protein.
SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gorilla gorilla gorilla (Lowland gorilla).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Gorilla.
                                                                                                                                            .,
                                                                                                        Length 352;
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 (BY SIMILARITY).
(BY SIMILARITY).
(BY SIMILARITY).
(BY SIMILARITY).
                                                     ULFATION (BY SIMILARITY)
F4E2F47135AF658A CRC64;
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                                                                                                          DB 1;
                                                                                                   72.1%; Score 1370; DB 1; 76.1%; Pred. No. 7.4e-74; live 30; Mismatches 46
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15 81
40527 MW;
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15-JUL-1998 (Rel. 36, Last seq
16-OCT-2001 (Rel. 40, Last anno
                                                                                                                                       Matches 261; Conservative
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P56439;
                                               MOD_RES
SEQUENCE
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EMBL; AF005659; AAB62553.1; -.
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_FI_1; 1.
PROSITE; PS500262; G_PROTEIN_RECEP_FI_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.

G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 YDIDXYTSEPCQKTNVKQIAARLLPPLXSLVFIFGFVGNMLVILLILINCKRLKSMTDIYL 69
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
C-C chamckine receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCR5).
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Papio anubis (Olive baboon).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                        Length 352;
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SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
DOBGFCB9FE5EAC84 CRC64;
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EXTRACELLULAR (POTENTIAL)
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CYTOPLASMIC (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
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ive 30; Mismatches
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NCBI_TaxID=9557, 9555;
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                                                                                                                                                                                                                                                                                                                                      Benton P.A., Timanus D.K., Shearer M.H., Lee D.R., Kennedy R.C.;
Submitted (NOV-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
                                                                                                                                                                                                                        Saksena N.K., Wang B., Novembre F.J., Bolton W., Smit T.K., Lal R.B.; "Species-specific changes in the CCR5 gene from African and Asian
                                                                                                                                                                                                                                                                                                                                                                                     MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL. MAY PLAY A ROLE
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PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBCELLULAR LOCATION: Integral membrane protein. SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                           SPECIES=P.hamadryas;
MEDLINE=97268687; PubMed=9108095;
Edinger A.L., Amedee A., Miller K., Doranz B.J., Endres M.,
Sharzon M., Samson M., Lu Z.H., Clements J.E., Murphey-Corb M.,
Peiper S.C., Parmentier M., Broder C.C., Doms R.W.;
"Differential utilization of CCR5 by macrophage and T cell tropic simian immunodeficiency virus strains.";
Proc. Natl. Acad. Sci. U.S.A. 94:4005-4010(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-LINKED (GLCNAC. . .) (POTENTIAL). 5E1504A9BA1FE8B2 CRC64;
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CYTOPLASMIC (POTENTIAL).
BY SIMILARITY.
SULFATION (BY SIMILARITY).
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CYTOPLASMIC (POTENTIAL).
2 (POTENTIAL).
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CYTOPLASMIC (POTENTIAL).
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                                                                                                                                                                                          SPECIES=P.hamadryas;
MEDLINE=99210133; PubMed=10195758;
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EMBL, AF105288, AAD20557.1;
EMBL, AF105289; AAD20558.1;
EMBL, AF105290; AAD20558.1;
EMBL, AF105290; AAD20559.1;
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352 AA;
                                                                                                                                                                                                                                                          nonhuman primates."
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MEDITIE=96291862; PubMed-8663314; Raport C.J., Gosling J., Schweichart V.L., Gray P.W., Charo I.F.; Molecular cloning and functional characterization of a novel human CC chemokine receptor (CCR5) for RANTES, MIP-lbeta, and MIP-lalpha.", J. Biol. Chem. 271:17161-17166(1996).
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                                                                                                                                                                                                                                                                                                                                                                                                                            142 IVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RG 197
                                                                                                                                                              82 LNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA 141
                                                                                 24 FDYDY--GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL 81
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01-0CT-1996 (Rel. 34, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine receptor type 5 (C-C CRR-5) (CC-CR-5) (CCR-5) (CHV-1 fusion co-receptor) (CHEMR13) (CD195 antigen).
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MEDLINE=96295970; PubMed=8699119;
Combadiere C., Ahuja S.K., Tiffany H.L., Murphy P.M.;
Cloning and functional expression of CC CKR5, a human monocyte CC chemokine receptor selective for MIP-1(alpha), MIP-1(beta), and
                                                                                                   198 WNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWT
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P51681; 014692; 014693; 014695; 014696; 014699; 014699; 014700; 014701; 014702; 014703; 014704; 014705; 014706; 014707; 014708; 015538; Q9UPA4;
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Length 352;
                                         45; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    318 VFFRKHITKRFCKQCPVFYRETVDGVTSTNTPSTGEQEVSAGL 360
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  72.0%; Score 1368; DB 1; 76.1%; Pred. No. 9.6e-74;
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                                       31; Mismatches
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Biochemistry 35:3362-3367(1996).
  Query Match 72.0%
Best Local Similarity 76.1%
Matches 261; Conservative
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Homo sapiens (Human).
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-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA, MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY INCREASING THE INTRACELLULAR CALCIUM IONS LEVEL. MAY PLAY A ROLE
                                                                                                                                                                                                                                                                                          Zhang L., Carruthers C.D., He T., Huang Y., Cao Y., Wang G., Hahn B.,
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE-98049523; PubMed=9388201;
Mummidl S., Ahuja S.S., McDaniel B.L., Ahuja S.K.;
Mummidl S. C chemokine receptor 5 (CCR5) gene. Multiple transcripts
With 5'-end heterogeneity, dual promoter usage, and evidence for
polymorphisms within the regulatory regions and noncoding exons.";
J. Biol. Chem. 272:30662-30671(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Farzan M., Mirzabekov T., Kolchinsky P., Wyatt R., Cayabyab M., Gerard N.P., Gerard C., Sodroski J., Choe H.;
"Tyrosine sulfation of the amino terminus of CCR5 facilitates HIV-1
                                                                                                                             MEDINE=98001387; PubMed=9343222; Kubat D.; Kabat D.; Kubmann S.E., Platt E.J., Kozak S.L., Kabat D.; Platt E.J., Rozak S.L., Kabat D.; Polymorphisms in the CCR5 genes of African green monkeys and mice implicate specific amino acids in infections by simian and human immunodeficiency viruses."; J. Virol. 71:8642-8656(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry into CD4+ cells is mediated by the chemokine receptor 5.";
Muzny D., Chen C.N., Evans C., Fitzgerald M., See L.H., Tang M., Porcel B.M., Dragan Y., Giacalone J., Pae A., Powell E., Solinsky K.A., Desilva U., Diaz-Perez S., Zhou X., Yu Y., Watanabe M., Doggett N., Garcia D., Sagripanti J.L.; Submitted (MAY-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                "HIV type 1 subtypes, coreceptor usage, and CCR5 polymorphism."; AIDS Res. Hum. Retroviruses 13:1357-1366(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CHARACTERIZATION OF ITS HIV-1 RECEPTOR FUNCTION.
MEDLINE-96260017; PubMed-8649511;
Deng H., Liu R., Ellmeder W., Choe S., Unutmaz D., Burkhart M.,
di Marzio P., Marmon S., Sutton R.E., Hill C.M., Davis C.B.,
Peiper S.C., Schall T.J., Littman D.R., Landau N.R.;
"Identification of a major co-receptor for primary isolates of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CHARACTERIZATION OF ITS HIV-1 RECEPTOR FUNCTION.
MEDILINE-90260018; PUDMed-8649512;
Dradic T., Litwin V., Allaway G.P., Martin S.R., Huang Y.,
Nagashima K.A., Cayanan C., Maddon P.J., Koup R.A., Moore J.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Magierowska M., Barre-Sinoussi F., Issafras H., Theodorou I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (MAR-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A., AND VARIANT ARG-178
                                                                                                                                                                                                                                                        SEQUENCE FROM N.A., AND POLYMORPHISMS.
MEDLINE=98022612; PubMed=9359654;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry.";
Cell 96:667-676(1999).
                                                                                                            SEQUENCE FROM N.A.
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SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation;
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CYTOPLASMIC (POTENTIAL).
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EMBL; U57840; AAC50598.1; -
EMBL; U57840; AAB17071.1; -
EMBL; U95326; AAB57793.1; -
EMBL; AF011500; AAB65700.1; -
EMBL; AF011500; AAB65700.1; -
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EMBL; AF011503; AAB65702.1; -
EMBL; AF011503; AAB65702.1; -
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EMBL; AF011533; AAB65733.1; -
EMBL; AF011533; AAB65733.1; -
EMBL; AF011534; AAB65731.1; -
EMBL; AF011535; AAB65731.1; -
EMBL; AF011537; AAB65731.1; -
EMBL; AF011535; AAB65731.1; -
EMBL; AF011535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL, AFO11519; ARB65719.1; EMBL, AFO11520; AAB65720.1; EMBL, AFO11521; AAB65721.1; EMBL, AFO11521; AAB65721.1; EMBL, AFO11523; AAB65723.1; EMBL, AFO11524; AAB65724.1; EMBL, AFO11525; AAB65724.1; EMBL, AFO11526; AAB65726.1; EMBL, AFO11529; AAB65729.1; EMBL, AFO11539; AAB65730.1; EMBL, AFO11531; AAB65731.1; EMBL, AFO11531; AAB65731.1; EMBL, AFO11531; AAB65733.1; EMBL, AFO11534; AAB65733.1; EMBL, AFO11535; AAB65735.1; EMBL, AFO11535; AAB65737.1; EMBL, AFO11535; AAB65737.1; EMBL, AFO11535; AAB65737.1; EMBL, AFO11535; AAB65737.1; EMBL, AFO11237; AAB65737.1; EMBL, AFO11237; AAB65737.1; EMBL, AFO11237; AAB94735.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00237; GPCRRHODOPSN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-99416438; PubMed-10486970;
Zhang Y.-P.;
Zhang Y.-W., Ryder O.A., Zhang Y.-P.;
Sequence evolution of the CCRS chemokine receptor gene in primates.";
Mol. Biol. Evol. 16:1145-1154(1999).
-!- FUNCTION: RECEPTOR FOR A C-C TYPE CHEMOKINE. BINDS TO MIP-1-ALPHA,
MIP-1-BETA AND RANTES AND SUBSEQUENTLY TRANSDUCES A SIGNAL BY
INCREASING THE INTRACELLULAR CALCIUM IONS LEWEL. MAY PLAY A ROLE.
IN THE CONTROL OF GRANULOCYTIC LINEAGE PROLIFERATION OR
                                                                                                                                                                                                                                                                                                       142 IVHAVFALKARTVTFGVVTSVITWLVAVFASVPGIIFTKCQKEDSVYVCGPYFP----RG 197
                                                                                                                                                                                                                                                                                                                                                            WNNFHTIMRNILGLVLPLLIMVICYSGILKTLLRCRNEKKRHRAVRVIFTIMIVYFLFWT 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PYNIVILLNIFQEFFGLSNCESTSQLDQATQVIETLGMTHCCINPIIYAFVGEKFRRYLS 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae; Colobinae;
Trachypithecus.
                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                      81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
C-C chemokine_receptor type 5 (C-C CKR-5) (CC-CKR-5) (CCR-5)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: Integral membrane protein. SIMILARITY: BELONGS TO FAMILY 1 OF G-PROTEIN COUPLED RECEPTORS.
                                                                                                                                                                                                                                   LNLAISDLLFLITLPLWAHSAANEWVFGNAMCKLFTGLYHIGYFGGIFFIILLTIDRYLA
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                                                                                                                                                                                Score 1364; DB 1; Length 352; Pred. No. 1.6e-73; 32; Mismatches 46; Indels
                                                                                                                            -> S (IN DBSNP:1800939)
EXTRACELLULAR (POTENTIAL)
           7 (POTENTIAL)..
CYTOPLASMIC (POTENTIAL).
BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -> H (IN INCCR5-72A).
                                                                                                   Y -> D (IN INCCR5-71A).
/FTId=VAR_003481.
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                                                                                          SULFATION
                                                   SULFATION
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les 259; Conservative
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097879;
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There are no restrictions on its ng as its content is in no way red. Usage by and for commercial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 FDYDY --GAPCHKFDVKQIGAQLLPPLYSLVFIFGFVGNMLVVLILINCKKLKCLTDIYL 81
                                                                                              PFam; PF00001; / Lul_1, ...
PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; 1.
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2; 1.
G-protein coupled receptor; Transmembrane; Glycoprotein; Sulfation.
30 EXTRACELLULAR (POTENTIAL).
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5 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
6 (POTENTIAL).
EXTRACELLULAR (POTENTIAL).
7 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
STIMLARITY.
SULFATION (BY SIMILARITY).
SULFATION (BY SIMILARITY).
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CYTOPLASMIC (POTENTIAL).
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EXTRACELLULAR (POTENTIAL).
3 (POTENTIAL).
CYTOPLASMIC (POTENTIAL).
4 (POTENTIAL).
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SULFATION (BY SIMILARITY)
4366F148D3A5938F CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Score 1363; DB 1;
Pred. No. 1.9e-73;
                                                                                                                                                                                                                                                                                                                                                                                                                                   ed. No. 1.9e
Mismatches
                                               or send an email to license@isb-sib.ch)
                                                                      EMBL; AF075443; AAD19855.1; -.
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
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75.88;
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